

Substitute Form PTO-1449 (Modified)	U.S. Department of Commerce Patent and Trademark Office	Attorney's Docket No. 10886-047001	Application No. 09/322,067
Information Disclosure Statement by Applicant (Use several sheets if necessary) (37 CFR §1.98(b))		Applicant	
		Filing Date May 27, 1999	Group Art Unit

U.S. Patent Documents

Examiner Initial	Desig. ID	Patent Number	Issue Date	Patentee	Class	Subclass	Filing Date If Appropriate
	AA						
	AB						
	AC						
	AD						
	AE						
	AF						
	AG						
	AH						
	AI						
	AJ						
	AK						

Foreign Patent Documents or Published Foreign Patent Applications

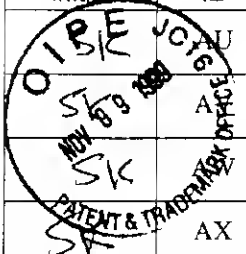
Examiner Initial	Desig. ID	Document Number	Publication Date	Country or Patent Office	Class	Subclass	Translation	
							Yes	No
	AL							
	AM							
	AN							
	AO							
	AP							

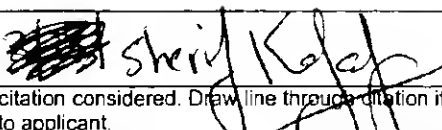
Other Documents (include Author, Title, Date, and Place of Publication)

Examiner Initial	Desig. ID	Document
SK	AQ	Wallace et al., "Derivation of 3D coordinate templates for searching structural databases: Application to Ser-His-Asp catalytic triads in the serine proteinases and lipases", Protein Science, 1996, Vol. 5, pages 1001-1013, see abstract
SK	AR	Escalier et al., "Pairwise and multiple identification of three-dimensional common substructures in proteins", Journal of Computational Biology, 1998, Vol. 5, No. 1, pages 41-56
SK	AS	Clarke et al., "Evidence from cassette mutagenesis for a structure function motif in a protein of unknown structure", Science, 22 April 1988, Vol. 240, pages 521-523
SK	AT	Viswanadhan et al., "New joint prediction algorithm (Q ₇ -Jasep) improves the prediction of protein secondary structure", Biochemistry, 1991, Vol. 30, pages 11164-11172

Examiner Signature <i>Sherif Kafafi</i>	Date Considered 3/15/2000
EXAMINER: Initials citation considered. Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant.	

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		Kalhammer et al., "Ras-binding domain: predicting function versus folding" FEBS Letters. 1997, Vol. 414, pages 599-602
		Chao et al., "Aligning two sequences within a specified diagonal band", CABIOS, 1992, Vol. 8, No. 5, pages 481-487
		Rychlewski et al., "Functional insight from structural predictions: Analysis of the Escherichia coli genome" March 1999, Vol. 8, pages 614-624
	AX	Pawlowski et al., "The Helicobacter pylori genome: From sequence analysis to structural and functional predictions", Proteins: Structure, Function and Genetics. July 1999, Vol. 36, pages 20-30
SK	AY	Zhang et al., "From fold predictions to function predictions: Automation of functional site conservation analysis for functional genome predictions", Protein Science. May 1999, Vol. 8, pages 1104-1115

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